OIPE

RAW SEQUENCE LISTING

DATE: 01/11/2002

PATENT APPLICATION: US/09/759,990

TIME: 13:19:48

Input Set : D:\31276-20026.txt

Output Set: N:\CRF3\01112002\I759990.raw

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ENTERED
 3 <110> APPLICANT: AntiCancer, Inc.
 4
         Xu, Mingxu
 5
         Han, Oinghong
   <120> TITLE OF INVENTION: HIGH EXPRESSION AND PRODUCTION OF HIGH
         SPECIFIC ACTIVITY RECOMBINANT S-ADENOSYLHOMOCYSTAINASE
         (SAHH) AND IMPROVED ASSAYS FOR S-ADENOSYLMETHIONINE (SAM)
12 <130> FILE REFERENCE: 31276-20026.00
14 <140> CURRENT APPLICATION NUMBER: US 09/759,990
15 <141> CURRENT FILING DATE: 2001-01-12
17 <150> PRIOR APPLICATION NUMBER: US 60/176,444
18 <151> PRIOR FILING DATE: 2000-01-14
20 <160> NUMBER OF SEQ ID NOS: 5
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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25 <211> LENGTH: 1461
26 <212> TYPE: DNA
27 <213> ORGANISM: Unknown
29 <220> FEATURE:
30 <223> OTHER INFORMATION: Nucleotide sequence encoding SAHH
32 <400> SEQUENCE: 1
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34 catgttctcg gccgtaagga acttaccctt gctgagaagg aaatgccagg tcttatggtt
                                                                        120
35 cttcgtgagc gttattccgc ttctaagcca ttgaagggtg tcagaatctc tggttccctc
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36 cacatgacag tecagacage egteeteate gagacaetea cagetettgg tgetgatgte
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37 agatgggett cetgeaacat ettetetaca caagatacag eegetgetge tategttgte
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38 ggcccaacag gcacaccaga gaagccagcc ggtatcccag tcttcgcctg gaagggcgaa
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39 acacteceag aataetggga gaacaeatae egegetetea eatggeeaga tggteaagge
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40 ccacagcagg ttgtcgatga tggtggtgat gctacactcc tcatctccaa gggcttcgaa
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41 ttcgaaacag ccggtgctgt tccagagcca acagaagctg acaacctcga ataccgctgc
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600
43 geoggeatga aeggtgttte egaagagaea acaacaggtg tecacegeet etaceagete
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44 gagaaggagg gcaaactcct cttcccagcc atcaacgtca acgacgctgt tacaaagtcc
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45 aagttegata aeatetaegg etgeegeeae teeettateg atggtateaa eegtgettee
                                                                        780
46 gatgtcatga tcggcggcaa gacagctctc gtcatgggtt acggcgatgt cggcaagggc
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47 tgcgctcaat ccctccgtgg ccaaggcgct cgcgttatca tcacagaact cgacccaatc
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48 tgcgctctcc aggctgccat ggaaggctac caggtccgcc gcatcgagga agtcgtcaag
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49 gatgtcgata tcttcgttac atgcacagga aactgcgata tcatctctgt tgacatgatg
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50 gcccagatga aggataaggc tattgtcggt aacatcggcc acttcgataa cgaaattgat
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51 acagatggee teatgaaata eecaggeate aageacatee caatcaagee agaataegae
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52 atgtgggaat teccagatgg ceaegetate etecttettg etgagggeeg cettettaae
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53 cttggctgcg ctacaggtca cccatctttc gttatgtcaa tgtcattcac aaaccagaca
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54 ctcgctcagc tcgacctcta cgaaaagaga ggaaatctcg agaagaaggt ttacacactt
                                                                       1320
55 ccgaagcate tegatgaaga agtegetege etecaceteg gatetetega tgtecacett
                                                                       1380
56 acaaagetta cacagaagca ggetgaetae atcaaegtte cagttgaggg teettacaag
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59 <210> SEQ ID NO: 2
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60 <211> LENGTH: 33

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101 ctacgaatgg caataattcc taggtacgta

30

VERIFICATION SUMMARY

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